

# Near-infrared Raman spectroscopy to detect anti-*Toxoplasma gondii* antibody in blood sera of domestic cats: quantitative analysis based on partial least-squares multivariate statistics

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**Abstract.** Toxoplasmosis is an important zoonosis in public health because domestic cats are the main agents responsible for the transmission of this disease in Brazil. We investigate a method for diagnosing toxoplasmosis based on Raman spectroscopy. Dispersive near-infrared Raman spectra are used to quantify anti-*Toxoplasma gondii* (IgG) antibodies in blood sera from domestic cats. An 830-nm laser is used for sample excitation, and a dispersive spectrometer is used to detect the Raman scattering. A serological test is performed in all serum samples by the enzyme-linked immunosorbent assay (ELISA) for validation. Raman spectra are taken from 59 blood serum samples and a quantification model is implemented based on partial least squares (PLS) to quantify the sample's serology by Raman spectra compared to the results provided by the ELISA test. Based on the serological values provided by the Raman/PLS model, diagnostic parameters such as sensitivity, specificity, accuracy, positive prediction values, and negative prediction values are calculated to discriminate negative from positive samples, obtaining 100, 80, 90, 83.3, and 100%, respectively. Raman spectroscopy, associated with the PLS, is promising as a serological assay for toxoplasmosis, enabling fast and sensitive diagnosis. © 2010 Society of Photo-Optical Instrumentation Engineers. [DOI: 10.1117/1.3463006]

**Keywords:** near-infrared Raman spectroscopy; multivariate statistical analysis; partial least squares; toxoplasmosis; cats; serology; indirect enzyme-linked immunosorbent assay.

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## 1 Introduction

Toxoplasmosis is a disease widely spread in many countries and it is caused by the *Toxoplasma gondii* protozoan. This disease presents a threat to the public health because the human infection by the protozoan is very common and can cause severe life-threatening diseases in immune compromised patients and in newborns with congenital toxoplasmosis. In addition *T. gondii* infection is important in veterinary medicine due to fetal abortions and neonatal losses of livestock.<sup>1</sup> Toxoplasmosis transmission occurs by ingestion of sporulated oocysts released with the feces of infected cats and other felids or by cysts present in the tissues of an infected animal. Humans can be contaminated via congenital infection, fecal matter residue consumption, or ingestion of uncooked meat.<sup>2</sup> Cats are the main cause for the toxoplasmosis transmission to hu-

mans and other animals.<sup>3</sup> The disease is found in several animal species that present important economic interest and it can cause severe profit losses if not detected.

Antibodies or immunoglobulins (Igs) are gamma globulin proteins that are found in blood or other body fluids of vertebrates, and they are used by the immune system to identify and neutralize foreign objects, such as bacteria and viruses. The diversity of antibodies enables the immune system to recognize an equally wide diversity of antigens. The unique part of the antigen recognized by an antibody is called an epitope. These epitopes bind with their antibodies in a highly specific interaction, called induced fit, that enables antibodies to identify and bind with only their unique antigen in the midst of the millions of different molecules that make up an organism. Specific IgM antibodies are the best indicators of a recent *T. gondii* infection in several species and can be detected from blood samples, indicating an early stage of the infection up to 12 weeks later, while IgG antibodies charac-

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terize a chronic disease and may be present up to 52 weeks after the infection took place.<sup>4–6</sup> Both antibodies can be used to detect several specific serum antigens with different infection levels as *Neospora caninum*, *Leishmania* spp, *Histomonas meleagridis*, and *Cryptosporidium parvum*, among others, and the antibody class investigation must be defined according to the study goal.<sup>7–10</sup>

Enzyme-linked immunosorbent assay (ELISA) is the most widely used serological method to diagnose toxoplasmosis in domestic animals, since it provides a good sensitivity to detect IgG or IgM for *T. gondii* protozoan.<sup>11</sup> Other serological assays in addition to ELISA are being used for IgG and IgM identification; for instance, chemiluminescence immunoassay (CLIA), enzyme-linked fluorescent assay (ELFA), and indirect immunofluorescent assay (IFA). All of these provide similar sensitivities, specificities, and positive and negative predictive values.<sup>12</sup> Nevertheless, these methods are not fast enough due to the necessity of sample preparation, including steps of washing, dilution, and incubation. Furthermore, the protein source used in the ELISA test usually contains multiple *T. gondii* antigens that may increase nonspecific cross-reactivity providing false positive results.<sup>13</sup> On the other hand, the use of other optical diagnosis techniques such as Fourier transform IR spectroscopy and near-IR Raman spectroscopy would enable a high sensitivity for molecular detection and they could be done faster and less expensively. Since they are nondestructive tests, there is no need for a sample preparation and, therefore, they can be automated.<sup>14</sup> Riley et al.<sup>15</sup> reported the use of the Fourier transform IR spectroscopy technique in conjunction to a partial least-squares (PLS) regression model to diagnose the failure of transfer of passive immunity in horses. They found sensitivity, specificity, accuracy, and predictive values for diagnosis as good as those obtained by using the radial immunodiffusion assay (RID).

The use of dispersive near-IR Raman spectroscopy has been expanding in the biomedical field.<sup>14,16,17</sup> This vibrational technique could help to determine the molecular composition of materials and is capable of a fast, accurate, and real time biological diagnosis.<sup>14</sup> Its use is being extended to the veterinary field.<sup>18,19</sup> Raman spectroscopy is a useful tool for quantitative biological analysis as well as for clinical diagnosis *in situ* and *ex situ*, which can be used to reduce the risk of invasive procedures.<sup>20–22</sup> It provides a high specificity to identify and quantify metabolites and pharmaceutical products<sup>23,24</sup> and it has the potential to identify and characterize microorganisms (bacteria and fungi) rapidly.<sup>25–27</sup> It can be used to determine the physiological conditions and the molecular composition of the tissues and organs through a noninvasive or minimally invasive method.<sup>21,28–31</sup> Torreggiani and Fini<sup>32</sup> studied the connection between the protein avidin and biotin through Raman spectroscopy. This study indicated that Raman spectroscopy could be used to detect specific antibodies due to its high sensitivity for molecular detection. Application of Raman spectroscopy to detect the presence of antibodies against *T. gondii* in cat blood serum<sup>18</sup> and in human colostrum has been reported.<sup>19</sup>

To be an effective and sensitive technique for biological material identification and classification, Raman spectroscopy requires an appropriate statistical method that can identify spectral differences that originate from differences in the sample composition that occur within the data set. Multivari-

ate statistics frequently used for quantitative analysis in Raman spectroscopy are the PLS regression and principal component regression (PCR). Such techniques can promote a quantitative analysis of compounds presented in the samples, even in the presence of multiple components where multicollinearity cannot be avoided. It was very successful in quantifying biologically relevant analytes such as glucose, ammonia, glutamine, glutamate, and lactate diluted in complex mixtures.<sup>33</sup> Berger et al.<sup>23</sup> determined the concentrations of the metabolites glucose, cholesterol, triglycerides, urea, total protein, and albumin in human blood serum and human total blood by Raman spectroscopy and PLS regression. Recent applications of IR spectroscopy associated with PLS regression include a device for bedside monitoring of subcutaneous interstitial glucose using near-IR spectrometry,<sup>34</sup> pulse glucometry for a noninvasive measurement of blood glucose,<sup>35</sup> and determination of glucose in clinically relevant concentrations in artificial aqueous humor with Raman spectroscopy.<sup>36</sup>

The objective of this work was to develop a model for blood sera assay based on Raman spectroscopy and multivariate statistical analysis PLS regression, for the quantitative analysis of *T. gondii* antibody (IgG) in the sera of domestic cats without and with diagnosed toxoplasmosis that have been submitted to the quantitative evaluation of IgG by an indirect ELISA test, and to develop a diagnostic model for discriminating positive from negative sera. The ELISA assay is widely used for serology and was used as a gold standard for detecting IgG in the blood sera included in the PLS model. A preliminary study showing the potential of the dispersive near-IR (NIR) Raman spectroscopy technique for detecting *T. gondii* antibodies (IgG) in cat blood sera was previously reported by our group,<sup>18</sup> indicating feasibility for such optical approach.

## 2 Materials and Methods

### 2.1 Domestic Cat Blood Serum Samples

A total of 59 cat blood serum samples were obtained from the Protection Animal Society (São Paulo, SP, Brazil) and used in the presented work. The samples were previously evaluated by the ELISA test, and following that, the Raman spectrum was taken for each sample.

### 2.2 ELISA

An ELISA test was performed at the Department of Pathology of Universidade Estadual Paulista (UNESP) (Jaboticabal, SP, Brazil) aiming to detect the presence of anti-*T. gondii* antibodies (IgG) in the cat serum. The ELISA test was performed following the protocol described by Domingues et al.<sup>37</sup> Serological results were quantified and defined as a set of ELISA levels (between 0 and 9), corresponding to a range of the ELISA optical density (OD) of the samples,<sup>38</sup> according to Table 1.

### 2.3 Raman Spectroscopy

A Raman spectrometer was used to collect the spectra; this instrument is described elsewhere.<sup>18,24</sup> Briefly, the excitation source was a Ti:sapphire laser (Spectra Physics, model 3900S) pumped by a 6-W multiline argon laser (Spectra Physics, model 2709S), tuned to 830 nm. The Raman scattered signal from the sample was collected and coupled into the spectrometer entrance slit by a set of lenses. The Rayleigh

**Table 1** OD by ELISA test, corresponding ELISA levels, and IgG serology.

ELISA OD Range	ELISA Level	IgG Serology	ELISA OD Range	ELISA Level	IgG Serology
0.000–0.221	0	N	0.736–0.993	5	MP
0.222–0.298	1	MP	0.994–1.340	6	MP
0.299–0.403	2	MP	1.341–1.809	7	SP
0.404–0.544	3	MP	1.810–2.443	8	SP
0.545–0.735	4	MP	2.444–3.298	9	SP

(N, negative; MP, medium positive; and SP, strong positive).

scattering from the sample was strongly attenuated in the collecting system by an holographic notch filter (Kaiser Optical Systems, model HSPF 830AR2.0). The optical detection system was composed of a high coupling spectrograph (Kaiser Optical Systems, Inc.,  $f/1.8i$ ) and a liquid-nitrogen-cooled, deep depletion CCD camera and controller (Princeton Instruments, model EEV 1024×256 and model ST130, respectively). To collect the Raman signal with an optimal resolution, the spectrometer entrance slit width was set<sup>39</sup> to 200  $\mu\text{m}$ , with a spectral resolution of about 10  $\text{cm}^{-1}$ . The CCD controller was connected to a personal computer that stored and processed the Raman spectra.

The quartz cuvette with the serum sample was positioned facing the spectrometer entrance slit. The pumping laser beam impinged on the cuvette, forming a right angle with respect to the signal-collecting optical axis. The serum samples were irradiated with 80-mW laser power and an acquisition time of 100 s for each sample.

The optical detection system (spectrometer and CCD) was calibrated in wave numbers using an organic compound, indene ( $\text{C}_9\text{H}_8$ ), which has a well-established Raman spectrum that presents intense and well-distributed bands covering the whole spectral region between 800 and 1800  $\text{cm}^{-1}$ .

#### 2.4 Preprocessing of NIR Raman Spectra

The contribution of the fluorescence emission to the Raman spectra was eliminated by fitting and subtracting a third-order polynomial from the gross spectrum. After calibration and background subtraction, the Raman spectra were filtered using the MATLAB<sup>®</sup> software to reduce the high-frequency noise (associated with the shot and readout noise). Cosmic ray spikes were manually removed. Once filtered, spectra were normalized by the most intense band and submitted to a multivariate statistical analysis, composed of the PLS regression method.

#### 2.5 Multivariate Statistical Analysis—PLS method

Samples were classified according to their IgG serology into two groups for PLS analysis: negative and positive samples. Then the 59 preprocessed spectra were separated randomly in two groups: 49 were used as the calibration dataset and the remaining 10 as the validation (prediction) data set, including positive and negative samples in the calibration and validation

groups. The PLS model was implemented by using the calibration set and the corresponding latent variables (LVs) were calculated from the first LV to the 10th LV.

To determine the number of LVs to be included in the PLS model that provides the minimum error predicted by the PLS compared to the ELISA OD values, we calculated the root mean square error of prediction (RMSEP) for each LV used in the validation data set as follows:

$$\text{RMSEP} = \left[ \frac{\sum_{i=1}^n (c_i - ca_i)^2}{n} \right]^{1/2},$$

where  $c_i$  is the OD value predicted by the PLS for the sample  $i$ ,  $ca_i$  is the OD value measured by ELISA (the gold-standard technique), and  $n$  is the number of samples. The minimum value of RMSEP will occur in the optimal LV, which can be used to predict future samples.

Likewise, the parameters sensitivity (Se), specificity (Sp), accuracy, positive predictive value (PPV), and negative predictive value (NPV) for the Raman/PLS model were calculated. Those parameters are defined as  $\text{Se} = \text{TP}/(\text{TP} + \text{FN})$ ,  $\text{Sp} = \text{TN}/(\text{TN} + \text{FP})$ ,  $\text{accuracy} = (\text{TP} + \text{TN})/(\text{TP} + \text{FP} + \text{TN} + \text{FN})$ ,  $\text{PPV} = \text{TP}/(\text{TP} + \text{FP})$ , and  $\text{NPV} = \text{TN}/(\text{FN} + \text{TN})$ , where TP, FP, TN, and FN are the numbers of true positives, false positives, true negatives, and false negatives, respectively.

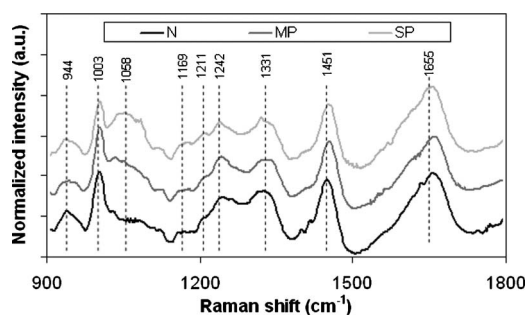
### 3 Results and Discussion

#### 3.1 ELISA

The 59 samples were classified as negative serum (N), medium positive (MP) and strong positive (SP) according to the following criterion (Table 1): N—ELISA level equal to 0, MP—ELISA levels ranging from 1 to 6, and SP—ELISA levels above 6. Based on this classification, it was obtained 13 N, 30 MP, and 16 SP blood sera.

#### 3.2 Raman Spectroscopy

Figure 1 shows the Raman spectra of three characteristic samples with N, MP, and SP ELISA serologies in the spectral range between 900 and 1800  $\text{cm}^{-1}$ . All spectra have a similar profile, with prominent band positions at 944, 1003, 1058, 1169, 1242, 1331, 1451, and 1655  $\text{cm}^{-1}$  for the three types of serology. An attempt to assign the bands presented in the serum, according to that reported by the literature, would be

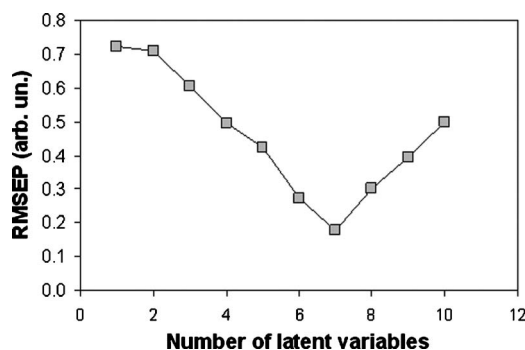


**Fig. 1** Raman spectra of cat blood sera with the following serology to toxoplasmosis (IgG): negative (N), medium positive (MP), and strong positive (SP).

band at  $944\text{ cm}^{-1}$  (proteins: C—C stretching mode of skeletal backbone  $\alpha$ -helix conformation); at  $1003\text{ cm}^{-1}$  (phenylalanine: symmetric ring breathing mode); at  $1058\text{ cm}^{-1}$  (phenylalanine and lipids: skeletal C—C stretch); at  $1169\text{ cm}^{-1}$  (amino acids and nucleic acids); at  $1242\text{ cm}^{-1}$  (amide III from proteins: N—H bending, C—N stretch;  $\text{CH}_2$  wag); at  $1331\text{ cm}^{-1}$  (proteins:  $\text{CH}_3$  and  $\text{CH}_2$  wagging; polynucleotide chain; phospholipids); at  $1451\text{ cm}^{-1}$  (lipids and proteins:  $\text{CH}_2$  deformation and proteins:  $\text{CH}_2$  bending); and at  $1655\text{ cm}^{-1}$  (lipids: C=C stretching and amide I from proteins: C=O stretching  $\alpha$ -helix conformation).<sup>14,16,40,41</sup> Differences observed in the band location compared to literature are within the spectrometer resolution ( $<10\text{ cm}^{-1}$ ). The overall spectrum profiles are very similar for negative and positive sera; however, it was possible to observe Raman bands with higher intensities for the MP and SP samples when compared to the N samples, which could be explained by the presence of IgG antibodies. For instance, the bands at  $1242\text{ cm}^{-1}$  (proteins) and  $1058\text{ cm}^{-1}$  (lipids) have significant differences, increasing with disease gravity.

### 3.3 PLS Model

The PLS regression model was used to develop a method to obtain the equivalent OD for the Raman spectra, thus obtaining an OD quantification and serology identification for the Raman/PLS model of cat blood sera. The data set was divided into two groups: 49 spectra for training the model and 10 spectra for validation (prediction), including negative and positive sera in both groups. First, we developed the calibration curve with the 49 spectra, in which the OD obtained by the ELISA test were used to determine the equivalent OD predicted by Raman/PLS. To determine how many LVs should be included in the PLS model to give the lowest prediction error, the 10 spectra from the validation data set were submitted to the PLS model using the number of LVs from 1 to 10. Then the RMSEP parameter was calculated for the OD values of the validation data set (Figure 2). We found that a lower RMSEP parameter is reached using seven LVs, so the OD values calculated using seven LVs for both training and validation groups versus the OD by ELISA are plotted in Fig. 3(a), whereas that for the validation group alone, separated by serology type, are shown in Fig. 3(b). We observed that all points lie close to the diagonal, zero-error line. The prediction RMSEP had a value of about  $\text{OD}=0.18$ , which is satisfactory

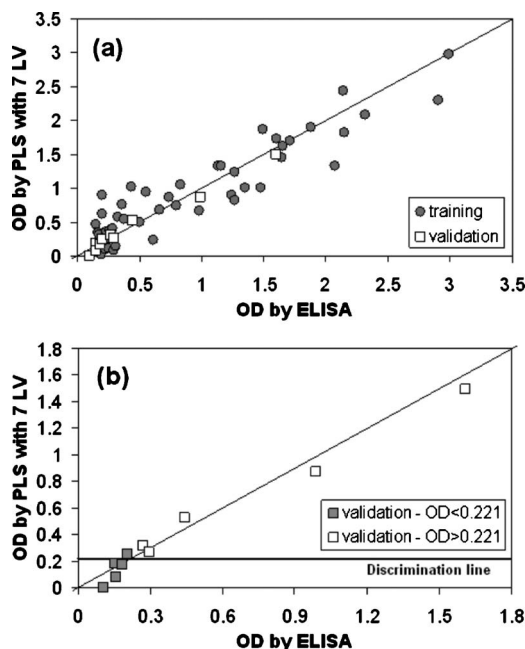


**Fig. 2** Plot of the RMSEP values of the OD predicted by the Raman/PLS model compared to OD values by ELISA using the first 10 LVs of the validation Raman data set.

for a method of rapid serology assay considering that the upper cutoff for the negative serology was  $\text{OD}=0.221$ .

### 3.4 Diagnosis Analysis

A comparison between OD values obtained by the Raman/PLS regression model and those obtained by the ELISA test in the validation data set is shown in Table 2. The table also shows the Se, Sp, accuracy, and PPV and NPV for the validation data set, considering positive the sera with  $\text{OD} > 0.221$ . We did not observe any false negative diagnosis, although we found only one sample out of the 10 analyzed that was identified as negative by the ELISA and diagnosed as positive by the Raman/PLS method. With respect to this



**Fig. 3** (a) Plot of the OD values predicted by the Raman-PLS model using seven LVs applied to cat blood sera versus the OD values obtained by ELISA test. Training group: 49 spectra, validation group: 10 spectra. The diagonal line represents the zero-error prediction line. (b) Plot of the OD achieved by the Raman/PLS model versus the OD from the ELISA test on a scale of 0 to 1.8 of the validation data set. The horizontal line represents  $\text{OD}=0.221$ , which discriminates negative from positive IgG serologies.



**Table 2** Classification capabilities of the Raman/PLS diagnosis correlated to the ELISA test (OD values) for the IgG from serum of domestic cats of the validation spectra.

Raman-PLS Diagnosis Validation data set	ELISA							
	OD≤0.221	OD>0.221	Total	Se (%)	Sp (%)	Accuracy (%)	PPV (%)	NPV (%)
ELISA OD≤0.221	4	0	4					
ELISA OD>0.221	1	5	6	100	80	90	83.3	100
Total	5	5	10					

Se, sensitivity; Sp, specificity; PPV, positive predictive value; NPV, negative predictive value.

sample, it was verified that this misclassified sample presented an OD value that is very close to the upper cutoff value for negative serum, and, therefore, the prediction seems to be more uncertain.

Riley et al. studied the diagnosis of the failure of transfer of passive immunity in horses using Fourier transform (FT) infrared spectroscopy employing PLS regression analysis.<sup>15</sup> They found parameter values of about 92, 96, 95, 86, and 98%, for Se, Sp, accuracy, PPV, and NPV, respectively. To compare Riley et al.'s work and the presented study, one should notice that the meaning of negative and positive is different in these works: FPT-positive in the Riley et al. paper is equivalent to serum-negative for toxoplasmosis in this study and vice versa. Therefore, the meanings of Se, Sp, PPV, and NPV are apposite in these works. These values are com-

parable to the ones presented in this model for toxoplasmosis diagnosis using Raman/PLS regression analysis (Table 2).

Table 3 compares the capabilities of the dispersive Raman spectroscopy in conjunction with a multivariate statistical analysis in diagnostic models for different diseases as reported in the literature.<sup>42-45</sup> We can see that our results on toxoplasmosis identification agree well with those obtained for other diseases using the same spectroscopic technique.

The NIR Raman technique has been proven to be suitable for biotissue analysis (including tissues, blood, and serum) due to the intrinsic vibrational information that is carried out by the Raman signal without interference from water. By using Raman spectroscopy, usually no sample preparation is required, a small amount of sample is used, the biochemical alterations could be obtained directly in real time, and de-

**Table 3** Comparison of the diagnostic capabilities of the Raman spectroscopy associated with multivariate statistical analysis in the diagnosis of several diseases.

Diagnostic Test for	Spectroscopic Method	Se (%)	Sp (%)
Oral carcinoma <sup>a</sup>	FT-Raman	100	55
Hepatitis C <sup>b</sup>	NIR dispersive Raman	92	88
Atherosclerosis in coronary artery <sup>c</sup>	NIR dispersive Raman		
Nonatherosclerotic plaque		78	93
Noncalcified plaque		90	83
Calcified plaque		100	100
Atherosclerosis in carotid artery <sup>d</sup>	FT-Raman		
Nonatherosclerotic plaque		96	93
Atherosclerotic plaque noncalcified		89	96
Atherosclerotic plaque calcified		100	100
Toxoplasmosis <sup>e</sup>	NIR dispersive Raman	100	80

<sup>a</sup>Oliveira et al.<sup>42</sup>

<sup>b</sup>Saade et al.<sup>43</sup>

<sup>c</sup>Silveira et al.<sup>44</sup>

<sup>d</sup>Nogueira et al.<sup>45</sup>

<sup>e</sup>This work

pending on the biochemical of interest, it results in a unique molecular fingerprint, which can be used to discriminate qualitatively and quantitatively the diseased tissue. Recent developments in Raman instrumentation include the use of low-cost diode lasers, CCD detectors with electronic cooling, fiber optic probes designed to reduce fiber background and increase SNR, and compact spectrometers with high throughput.<sup>46</sup>

## 4 Conclusions

Results indicated that the dispersive NIR Raman spectroscopy method in conjunction with a PLS regression analysis (Raman/PLS model) is effective in detecting toxoplasmosis in blood sera of domestic cats compared to the ELISA test. This method was shown to be very promising as an alternative technique for toxoplasmosis diagnosis, providing several advantages over conventional serological methods of diagnosis.

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